



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/500,218
Source: PT/10
Date Processed by STIC: 7/6/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
U.S. Patent and Trademark Office, 220 20th Street S.; Customer Window; Mail Stop Sequence, Crystal Plaza Two, Lobby Room 1B03, Arlington, VA 22202

Revised 05/17/04



PCT

RAW SEQUENCE LISTING

DATE: 07/07/2004

PATENT APPLICATION: US/10/500,218

TIME: 07:58:40

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\07062004\J500218.raw

3 <110> APPLICANT: Yoshihiro HORIUCHI
 5 <120> TITLE OF INVENTION: Hydroxamic acid derivative and MMP Inhibitor containing said derivative as
 6 active ingredient
 8 <130> FILE REFERENCE: 2004-0979A/WMC/001177
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/500,218
 11 <141> CURRENT FILING DATE: 2004-06-25
 13 <150> PRIOR APPLICATION NUMBER: PCT/JP02/13580
 14 <151> PRIOR FILING DATE: 2002-12-26
 16 <150> PRIOR APPLICATION NUMBER: JP 2001-397638
 17 <151> PRIOR FILING DATE: 2001-12-27
 19 <160> NUMBER OF SEQ ID NOS: 3
 21 <210> SEQ ID NO: 1
 22 <211> LENGTH: 44
 23 <212> TYPE: DNA
 24 <213> ORGANISM: Artificial Sequence *see p.2 for explanation*
 W--> 26 <220> FEATURE:
 W--> 26 <223> OTHER INFORMATION:
 W--> 26 <400> 1
 27 ggaattccat atgctgccgc tgccgagtg tggtgatgaa gatg 44
 29 <210> SEQ ID NO: 2
 30 <211> LENGTH: 51
 31 <212> TYPE: DNA
 32 <213> ORGANISM: Artificial Sequence *p.2*
 W--> 34 <220> FEATURE:
 W--> 34 <223> OTHER INFORMATION:
 W--> 34 <400> 2
 35 tttggatcct tagccgtaca ggctttgaat accttgatca tcgtcatcag g 51
 37 <210> SEQ ID NO: 3
 38 <211> LENGTH: 6
 39 <212> TYPE: PRT
 40 <213> ORGANISM: Artificial Sequence *p.2*
 42 <220> FEATURE:
 W--> 43 <221> NAME/KEY: Residue
 44 <222> LOCATION: (1)
 45 <223> OTHER INFORMATION: Xaa = 7-methoxycoumalin-4-yl proline
 47 <220> FEATURE:
 W--> 48 <221> NAME/KEY: Residue
 49 <222> LOCATION: (5)
 50 <223> OTHER INFORMATION: X = L-[N-(2,4-dinitrophenyl)-L-2,3-diaminopropionyl]-alanine.
 52 <400> SEQUENCE: 3
 WOK 53 Xaa Leu Gly Leu Xaa Arg
 54 1 5

pp 1-2
 Does Not Comply
 Corrected Diskette Needed

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/500,218

DATE: 07/07/2004
TIME: 07:58:41

Input Set : A:\PTO.DA.txt
Output Set: N:\CRF4\07062004\J500218.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; Xaa Pos. 1,5

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 5

Use of <220> Feature (NEW RULES):

error explanation
Sequence(s) are missing the <220> Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32) (Sec.1.823 of new Rules)

Seq#:1,2,3

VERIFICATION SUMMARY

DATE: 07/07/2004

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Input Set : A:\PTO.DA.txt

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L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:26 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:1, <213>
ORGANISM:Artificial Sequence
L:26 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:1, <213>
ORGANISM:Artificial Sequence
L:26 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:26
L:34 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:2, <213>
ORGANISM:Artificial Sequence
L:34 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:2, <213>
ORGANISM:Artificial Sequence
L:34 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:2,Line#:34
L:43 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3
L:48 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:3
L:53 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0